- (ii) MOLECULE TYPE: peptide
- (v) FRAGMENT TYPE: N-terminal
- (vi) ORIGINAL SOURCE:
 - (A) ORGANISM: alkalophilic Bacillus sp.
 - (B) STRAIN: 38-2
- (vii) IMMEDIATE SOURCE:
 - (B) CLONE: beta-cyclodextrin
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

Ala Pro Asp Thr Ser Val Ser Asn Lys Gln Asn Phe Ser Thr Asp Val

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REMARKS

The present Amendment is in response to Examiner's Supplemental Notice to Comply With Requirements For Patent Applications Containing Nucleotide Sequence And/Or Amino Acid Sequence Disclosures mailed June 2, 1993.

Applicants submit this Amendment to complete their disclosure of the sequences employed in the instant application pursuant to 37 C.F.R. § 1.821 et seq. Applicants also submit herewith a paper copy of the Sequence Listing and a floppy disk containing the Sequence Listing in computer readable form in compliance with 37 C.F.R. §§ 1.821 (c) and (e).

Examiner will note that Sequence 7 does contain 709 base pairs. The error in the amended sequence listing submitted May 3, 1993 was not that Sequence 7 has 677 base pairs but rather that the label "709" was

not positioned in the far right margin and hence was not noticed by the Examiner. This inadvertant error has been remedied by the present Amendment and the newly submitted sequence listing.

Entry of the present amendment is proper since no new matter has been introduced by the instant amendment to the Specification or by these submissions and since the content of the amendment to the Specification and the paper and computer readable copies of the Sequence Listing are the same.

Respectfully submitted,

LIMBACH & LIMBACH

Dated: 1/we 10,183

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Attorneys for Applicant